

# Python for Biologists - Local Development Setup

## Mac & Linux Installation Guide






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### Overview

This guide will walk you through setting up a professional Python development environment on your Mac or Linux machine. Follow these steps in order!

#### What you'll install:

-  Warp Terminal (modern terminal with AI features)
-  Git (for downloading course materials)
-  UV (Python package manager)
-  Cursor IDE (code editor with AI assistance)
-  Course notebooks repository

**Estimated Time:** 30-45 minutes

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### Step 1: Install Warp Terminal

Warp is a modern terminal with AI features that makes command-line work much easier for beginners.

#### Mac Users

1. Visit [warp.dev](https://warp.dev)
2. Click "Download for Mac"
3. Open the downloaded .dmg file
4. Drag Warp to your Applications folder
5. Open Warp from Applications
6. Sign in with your email (free account)

## Linux Users

1. Visit [warp.dev](https://warp.dev)
2. Download for Linux
3. Follow the installation instructions for your distribution
4. Launch Warp

**Alternative:** If you prefer, you can use your system's default terminal (Terminal on Mac, GNOME Terminal on Linux). All commands work the same!

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## Step 2: Check if Git is Installed

Before installing anything, let's check if you already have Git.

### Open Warp (or your terminal) and run:

```
git --version
```

If you see a version number (like `git version 2.39.0`):

-  **You're good to go!** Skip to Step 3.

If you see “command not found”:

- Continue with the installation instructions below.

## Mac Users: Install Git via Homebrew

### Option 1: Homebrew (Recommended)

First, install Homebrew (a package manager for Mac):

```
/bin/bash -c "$(curl -fsSL https://raw.githubusercontent.com/Homebrew/install/HEAD/install.sh)"
```

Follow the on-screen instructions and enter your password when prompted.

Then install Git:

```
brew install git
```

### Option 2: Xcode Command Line Tools

If you run `git --version` and get a popup asking to install developer tools, click “Install”. This is Apple’s built-in option (but Homebrew is better long-term).

## Linux Users: Install Git

### Ubuntu/Debian:

```
sudo apt update
sudo apt install git -y
```

### **Fedora:**

```
sudo dnf install git -y
```

### **Verify Git Installation**

```
git --version
```

You should now see a version number!

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## **Step 3: Clone the Course Repository**

Now let's download all the course materials from GitHub.

### **Navigate to where you want to store your course files:**

```
cd Documents
mkdir biology-python
cd biology-python
```

### **Clone the repository:**

```
git clone https://github.com/HocheggerLab/y3-bio-python.git
```

You'll see output like:

```
Cloning into 'y3-bio-python'...
remote: Enumerating objects...
Receiving objects: 100% done.
```

### **Navigate into the repository:**

```
cd y3-bio-python
```

✅ **Success!** You now have all the course notebooks on your computer.

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## **Step 4: Install UV (Python Package Manager)**

UV is a modern, fast Python package manager that will handle all your dependencies.

## Install UV:

### Mac & Linux:

```
curl -LsSf https://astral.sh/uv/install.sh | sh
```

## Restart your terminal

Close Warp and open it again (or open a new terminal tab).

## Verify installation:

```
uv --version
```

You should see the UV version number!

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## Step 5: Set Up Python Environment

Now let's install all the course packages (numpy, pandas, matplotlib, jupyter, etc.)

### Make sure you're in the course repository folder:

```
cd ~/Documents/biology-python/y3-bio-python
```

(Adjust the path if you put it somewhere else)

### Sync all course packages:

```
uv sync
```

This will:

- Read the `pyproject.toml` file
- Create a virtual environment (`.venv` folder)
- Install Python 3.12 and all course packages

This takes 2-3 minutes. You'll see output like:

```
Resolved 23 packages in 450ms  
Installed 23 packages in 2.1s
```

✅ **Done!** Your Python environment is ready.

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## Step 6: Download and Install Cursor IDE

Cursor is like VS Code but with AI superpowers - perfect for learning Python!

## Download Cursor:

1. Visit [cursor.com](https://cursor.com)
2. Click “Download for Mac” or “Download for Linux”
3. **Mac:** Open the .dmg file and drag Cursor to Applications
4. **Linux:** Follow the installation instructions for your distribution

## Launch Cursor:

1. Open Cursor from Applications (Mac) or your app menu (Linux)
2. Grant any permissions if prompted

## Sign in (optional but recommended):

- Click “Sign In” in the bottom-left
  - Create a free account or sign in with GitHub
  - This enables AI features
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# Step 7: Open the Repository in Cursor

## Option 1: From Terminal

In your terminal (make sure you’re in the y3-bio-python folder):

```
cursor .
```

This opens the current folder in Cursor!

## Option 2: From Cursor

1. Open Cursor
  2. File → Open Folder
  3. Navigate to ~/Documents/biology-python/y3-bio-python
  4. Click “Open”
- 

# Step 8: Install Required Cursor Extensions

You need two extensions to work with Jupyter notebooks.

## Install Python Extension:

1. Click the Extensions icon in the left sidebar (looks like building blocks)
  - Or press `Cmd+Shift+X`
2. Search for “Python”
3. Install the one by **Microsoft** (it has the most downloads)

## Install Jupyter Extension:

1. In the Extensions sidebar, search for “Jupyter”
2. Install the one by **Microsoft**
3. **Restart Cursor** after installation

**Note:** Cursor usually prompts you to install these automatically when you open a .ipynb file!

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## Step 9: Select Python Interpreter

Tell Cursor to use your virtual environment.

### Steps:

1. Press `Cmd+Shift+P` to open the command palette
2. Type “Python: Select Interpreter”
3. Select the interpreter that shows `.venv` in the path
  - Should look like: Python 3.12.x ( '.venv': venv )

**You only need to do this once per project!**

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## Step 10: Test Your Setup

Let's make sure everything works by running a notebook!

### Open a notebook:

1. In the Cursor file explorer (left sidebar), navigate to:  
`notebooks/lecture_1/`
2. Click on `L1_N1_collab_notebooks.ipynb`

### Run the first cell:

1. Look for the first code cell in the notebook
2. Click the  play button next to it
  - Or press `Shift+Enter`
3. Wait a moment for the kernel to start

### What should happen:





- The cell runs
- You see output below the cell
- A number appears in brackets like `[1]` next to the cell

## If it asks you to select a kernel:

1. Click “Select Kernel” at the top-right of the notebook
  2. Choose “Python Environments...”
  3. Select the one with `.venv` in the name
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## You're All Set!

Congratulations! Your development environment is fully set up. You can now:

-  Open any notebook in the notebooks/ folder
  -  Run cells with Shift+Enter
  -  Ask Cursor AI for help (press Cmd+L)
  -  Work offline with all course materials
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## Quick Reference Commands

```
# Navigate to your project
cd ~/Documents/biology-python/y3-bio-python
```

```
# Open in Cursor
cursor .
```

```
# Get latest course updates
git pull
```

```
# Add new Python packages (if needed)
uv add package-name
```

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## Troubleshooting

### “git: command not found”

- Make sure you completed Step 2
- On Mac, if Homebrew is installed: `brew install git`

### “uv: command not found”

- Restart your terminal after installing UV
- Make sure the install script completed successfully

### Notebook won't run / “No kernel”

- Make sure you selected the `.venv` interpreter (Step 9)

- Try restarting Cursor
- Make sure both Python and Jupyter extensions are installed

## “Cannot find module ‘numpy’” or similar

- Make sure you ran `uv sync` in Step 5
- Make sure you selected the correct interpreter (should show `.venv`)

## Extension installation fails

- Check your internet connection
  - Try restarting Cursor and installing again
  - Extensions sidebar: click the “...” menu → “Check for Extension Updates”
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## Getting Help

- **Cursor AI:** Press `Cmd+L` and ask questions about your code
  - **Course Seminars:** Bring setup questions to practical sessions
  - **Course Forums:** Post questions - other students can help!
  - **Instructor:** Email [hh65@sussex.ac.uk](mailto:hh65@sussex.ac.uk) for technical issues
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## What’s Next?

1. **Explore the notebooks:** Start with `notebooks/lecture_1/`
2. **Practice running cells:** Get comfortable with the interface
3. **Try Cursor AI:** Press `Cmd+L` and ask it to explain Python concepts
4. **Attend seminars:** We’ll work through exercises together!

**Remember:** Focus on learning Python, not mastering the IDE. The tools are here to help you code!

Happy coding! 🐍🔍